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21
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1 caacaatcgggatataaacc.....tattaaatcttgcarctgcr 7582
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description		Abl61744 Colon ade Abn97929 Human ret	Ads30988 Human gen Ad£59718 Human con	Aas84210 DNA encod		-	Adb72620 Human PAP Adc85361 Mouse Pap	Human	Adh58564 Human Na+	Abd32817 Human can	Aas67609 DNA encod	Ade09587 Novel DNA	Aad24195 Human syn	Aax77526 Human sec	Aaz59468 Human sec
SUMMARIES	AAA59215 AAX25665	ABL61744 ABN97929	ADF59718	AAS84210	AAS68626	ADA02882	ADB72620 ADC85361	ADM74477	ADH58564	ABD32817	AAS67609	ADE09587	AAD24195	AAX77526	AAZ59468
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* Query Match	98.8 98.8	85.0 84.9	67.4	49.3	43.7	40.2	40.2	40.2	38.9	38.9	38.7	38.4	38.1	38.1	38.1
Score	7489.6	6448	6124.6 5108	3738.4	3309.6	3049.8	3049.8	3049.8	2948	2948	2932	2908.2	2889.6	2888.6	2888.6
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PN JP 2002512530-A 11 23-APR-2002;
PN JP 2002512330-A)11
PD 23-APR-2002
PP 05-JUL-1998 JP 1999508244
PR 07-JUL-1998 JF 97/08815
PR 07-JUL-1997 FR 97/08815
PR FEEDERIC BESEME, JEAN LUC BLOND, OLIVIER BOUTON, BERNARD
                                                                                                                                                                                                                                                                                                                                                                                                              Endogenetic retroviral sequences, associated with autoimmune diseases and/or with pregnancy disorders.
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Strandedness: Single;
CC Topology: Linear;
CC Endogenetic retroviral sequences, associated with aut
CC Endogenetic retroviral sequences
CC and/or with pregnancy disorders
CC and/or with pregnancy disorders
FH Key Location/Qualifiers
FT source / Organism='Unidentified'.
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                                                                                                                                                         Description
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Genoscope.

Direct Submission

Submitted (20-JUL-2004) Genoscope - Centre National de Sequencage :

BP 191 91006 EVRY cedex - FRANCE (E-mail : seqrefégenoscope.cns.fr
- Web : www.genoscope.cns.fr)

1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies, a division of Invitrogen.

Location/Qualifiers
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                          BX368078 BX36707
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Li, W.B., Gruber, C., Jessee, J. and Polayes, D.
Full-length cDNA libraries and normalization
Unpublished
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full-length cDNA clone CS0DE012YJ24 of Placenta of Homo sapiens
(human).
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http://fulllength.invitrogen.com/ InVitroGen Corporation 1600
Faraday Avenue
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HTC; CNSLT_CDNA.
HOMO SAPIENS (human)
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